

SEQUENCE LISTING

<110> Cohen, Stanley N.
Li, Limin

<120> MAMMALIAN TUMOR SUSCEPTIBILITY GENES AND
THEIR USES

<130> FUNC-0027-CO5

<140> US 10/697,720

<141> 2003-10-29

<150> US 09/804,690

<151> 2001-03-12

<150> US 09/146,187

<151> 1998-09-01

<150> US 08/977,818

<151> 1997-11-25

<150> US 08/670,274

<151> 1996-06-13

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<151> 1996-01-16

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<151> 1995-11-16

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1 5 10 15

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aat gtc atc gct atg tac aaa gat ctc aaa cct gta ttg gat tca tat 156
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gtt ttt aat gat ggc agt tcc agg gag ctg gtg aac ctc act ggt aca 204
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atc cca gtg cgt tat cga ggt aat ata tat aat att cca ata tgc ctg 252
Ile Pro Val Arg Tyr Arg Gly Asn Ile Tyr Asn Ile Pro Ile Cys Leu
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Pro	Thr	Ser	Ser	Met	Thr	Ile	Lys	Thr	Gly	Lys	His	Val	Asp	Ala	Asn	
				85					90					95		
ggg	aaa	atc	tac	cta	cct	tat	cta	cat	gac	tgg	aaa	cat	cca	cgg	tca	396
Gly	Lys	Ile	Tyr	Leu	Pro	Tyr	Leu	His	Asp	Trp	Lys	His	Pro	Arg	Ser	
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Glu	Leu	Leu	Glu	Leu	Ile	Gln	Ile	Met	Ile	Val	Ile	Phe	Gly	Glu	Glu	
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Pro	Pro	Val	Phe	Ser	Arg	Pro	Thr	Val	Ser	Ala	Ser	Tyr	Pro	Pro	Tyr	
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Thr	Ala	Thr	Gly	Pro	Pro	Asn	Thr	Ser	Tyr	Met	Pro	Gly	Met	Pro	Ser	
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Pro	Gly	Cys	Pro	Tyr	Pro	Pro	Ala	Gly	Pro	Tyr	Pro	Ala	Thr	Thr	Ser	
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Gln	Ala	Glu	Leu	Asn	Ala	Leu	Lys	Arg	Thr	Glu	Glu	Asp	Leu	Lys	Lys	
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Ala	Glu	Val	Asp	Lys	Asn	Ile	Glu	Leu	Leu	Lys	Lys	Lys	Asp	Glu	Glu	
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Leu	Ser	Ser	Ala	Leu	Glu	Lys	Met	Glu	Asn	Gln	Ser	Glu	Asn	Asn	Asp	
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Gly Glu Ala Leu Arg Arg Gly Val Ile Asp Leu Asp Val Phe Leu Lys	
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cac gtc cgc ctc ctg tcc cgt aaa cag ttc cag cta agg gca cta atg	1164
His Val Arg Leu Leu Ser Arg Lys Gln Phe Gln Leu Arg Ala Leu Met	
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caa aag gca agg aag act gcg ggc ctt agt gac ctc tac tgacatgtgc	1213
Gln Lys Ala Arg Lys Thr Ala Gly Leu Ser Asp Leu Tyr	
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tatatatttct gtttcctttg ggtaaaaact ggcttttatt aatgcacttt ctaccctctg	1393
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<223> TSG101 nucleotide

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Ile Pro Val Arg Tyr Arg Gly Asn Ile Tyr Asn Ile Pro Ile Cys Leu	
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Trp Leu Leu Asp Thr Tyr Pro Tyr Asn Pro Pro Ile Cys Phe Val Lys	
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Pro Thr Ser Ser Met Thr Ile Lys Thr Gly Lys His Val Asp Ala Asn	
85 90 95	
Gly Lys Ile Tyr Leu Pro Tyr Leu His Asp Trp Lys His Pro Arg Ser	
100 105 110	
Glu Leu Leu Glu Leu Ile Gln Ile Met Ile Val Ile Phe Gly Glu Glu	
115 120 125	
Pro Pro Val Phe Ser Arg Pro Thr Val Ser Ala Ser Tyr Pro Pro Tyr	
130 135 140	
Thr Ala Thr Gly Pro Pro Asn Thr Ser Tyr Met Pro Gly Met Pro Ser	
145 150 155 160	
Gly Ile Ser Ala Tyr Pro Ser Gly Tyr Pro Pro Asn Pro Ser Gly Tyr	
165 170 175	
Pro Gly Cys Pro Tyr Pro Pro Ala Gly Pro Tyr Pro Ala Thr Thr Ser	
180 185 190	
Ser Gln Tyr Pro Ser Gln Pro Pro Val Thr Thr Val Gly Pro Ser Arg	
195 200 205	
Asp Gly Thr Ile Ser Glu Asp Thr Ile Arg Ala Ser Leu Ile Ser Ala	
210 215 220	
Val Ser Asp Lys Leu Arg Trp Arg Met Lys Glu Glu Met Asp Gly Ala	
225 230 235 240	
Gln Ala Glu Leu Asn Ala Leu Lys Arg Thr Glu Glu Asp Leu Lys Lys	
245 250 255	
Gly His Gln Lys Leu Glu Glu Met Val Thr Arg Leu Asp Gln Glu Val	

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Asn	Leu	Tyr	Ala	Glu	Glu	Asn	Ala	Ile	Glu	Asp	Thr	Ile	Phe	Tyr	Leu
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Pro	Val	Thr	Thr	Val	Gly	Pro	Ser	Arg	Asp	Gly	Thr	Ile	Ser	Glu	Asp
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Met	Val	Thr	Arg	Leu	Asp	Gln	Glu	Val	Ala	Glu	Val	Asp	Lys	Asn	Ile
Glu	Leu	Lys	Lys	Lys	Asp	Glu	Glu	Leu	Ser	Ser	Ala	Leu	Glu	Lys	
Met	Glu	Asn	Gln	Ser	Glu	Asn	Asn	Asp	Ile	Asp	Glu	Val	Ile	Ile	Pro
Thr	Ala	Pro	Leu	Tyr	Lys	Gln	Ile	Leu	Asn	Leu	Tyr	Ala	Glu	Glu	Asn
Ala	Ile	Glu	Asp	Thr	Ile	Phe	Tyr	Leu	Gly	Glu	Ala	Leu	Arg	Arg	Gly
Val	Ile	Asp	Leu	Asp	Val	Phe	Leu	Lys	His	Val	Arg	Leu	Leu	Ser	Arg
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35 40 45
Ile Pro Val Pro Tyr Arg Gly Asn Thr Tyr Asn Ile Pro Ile Cys Leu
50 55 60
Trp Leu Leu Asp Thr Tyr Pro Tyr Asn Pro Pro Ile Cys Phe Val Lys
65 70 75 80
Pro Thr Ser Ser Met Thr Ile Lys Thr Gly Lys His Val Asp Ala Asn
85 90 95
Gly Lys Ile Tyr Leu Pro Tyr Leu His Glu Trp Lys His Pro Gln Ser
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Asp Leu Leu Gly Leu Ile Gln Val Met Ile Val Val Phe Gly Asp Glu
115 120 125
Pro Pro Val Phe Ser Arg Pro Ile Ser Ala Ser Tyr Pro Pro Tyr Gln

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Ile	Ser	Pro	Tyr	Pro	Ser	Gly	Tyr	Pro	Pro	Asn	Pro	Ser	Gly	Tyr	Pro
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Gly	Cys	Pro	Tyr	Pro	Pro	Gly	Gly	Pro	Tyr	Pro	Ala	Thr	Thr	Ser	Ser
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Gln	Tyr	Pro	Ser	Gln	Pro	Pro	Val	Thr	Thr	Val	Gly	Pro	Ser	Arg	Asp
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Gly	Thr	Ile	Ser	Glu	Asp	Thr	Ile	Arg	Ala	Ser	Leu	Ile	Ser	Ala	Val
	210					215					220				
Ser	Asp	Lys	Leu	Arg	Trp	Arg	Met	Lys	Glu	Glu	Met	Asp	Arg	Ala	Gln
225					230					235					240
Ala	Glu	Leu	Asn	Ala	Leu	Lys	Arg	Thr	Glu	Glu	Asp	Leu	Lys	Lys	Gly
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His	Gln	Lys	Leu	Glu	Glu	Met	Val	Thr	Arg	Leu	Asp	Gln	Glu	Val	Ala
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Glu	Val	Asp	Lys	Asn	Ile	Glu	Leu	Leu	Lys	Lys	Lys	Asp	Glu	Glu	Leu
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Ser	Ser	Ala	Leu	Glu	Lys	Met	Glu	Asn	Gln	Ser	Glu	Asn	Asn	Asp	Ile
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Asp	Glu	Val	Ile	Ile	Pro	Thr	Ala	Pro	Leu	Tyr	Lys	Gln	Ile	Leu	Asn
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Glu	Ala	Leu	Arg	Arg	Gly	Val	Ile	Asp	Leu	Asp	Val	Phe	Leu	Lys	His
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Lys	Ala	Arg	Lys	Thr	Ala	Gly	Leu	Ser	Asp	Leu	Tyr				
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